



SEQUENCE LISTING

<110> Lawrence, Papsidero
Lyn, Dyster
Jana, Frustaci

<120> Detection and Treatment of Breast Cancer

<130> 3380/1I127-US4

<140> 09/834,794

<141> 2001-04-13

<150> 09/146,580

<151> 1998-09-03

<150> 60/071,899

<151> 1998-01-20

<150> 60/092,155

<151> 1998-07-09

<160> 35

<170> PatentIn version 3.0

<210> 1

<211> 127

<212> PRT

<213> Homo sapiens

<220>

<221> UNSURE

<222> (70)..(70)

<223> Xaa at position 70 is either Arg or Gly

<220>

<221> UNSURE

<222> (91)..(91)

<223> Xaa at position 91 is either Lys or Asn

<400> 1

Met Gln Gln Arg Gly Leu Ala Ile Val Ala Leu Ala Val Cys Ala Ala
1 5 10 15

Leu His Ala Ser Glu Ala Ile Leu Pro Ile Ala Ser Ser Cys Cys Thr
20 25 30

#3

Glu Val Ser His His Ile Ser Arg Arg Leu Leu Glu Arg Val Asn Met
35 40 45

Seq ID #3
Cys Arg Ile Gln Arg Ala Asp Gly Asp Cys Asp Leu Ala Ala Val Ile
50 55 60

Leu His Val Lys Arg Xaa Arg Ile Cys Val Ser Pro His Asn His Thr
65 70 75 80

Val Lys Gln Trp Met Lys Val Gln Ala Ala Xaa Lys Asn Gly Lys Gly
85 90 95

Asn Val Cys His Arg Lys Lys His His Gly Lys Arg Asn Ser Asn Arg
100 105 110

Ala His Gln Gly Lys His Glu Thr Tyr Gly His Lys Thr Pro Tyr
115 120 125

<210> 2

<211> 104

<212> PRT

<213> Homo sapiens

<220>

<221> UNSURE

<222> (47)..(47)

<223> Xaa at position 47 is either Arg or Gly

<220>

<221> UNSURE

<222> (68)..(68)

<223> Xaa at position 68 is either Lys or Asn

<400> 2

Leu Pro Ile Ala Ser Ser Cys Cys Thr Glu Val Ser His His Ile Ser
1 5 10 15

Arg Arg Leu Leu Glu Arg Val Asn Met Cys Arg Ile Gln Arg Ala Asp
20 25 30

Gly Asp Cys Asp Leu Ala Ala Val Ile Leu His Val Lys Arg Xaa Arg
35 40 45

Ile Cys Val Ser Pro His Asn His Thr Val Lys Gln Trp Met Lys Val
50 55 60

Gln Ala Ala Xaa Lys Asn Gly Lys Gly Asn Val Cys His Arg Lys Lys

65		70		75		80									
His	His	Gly	Lys	Arg	Asn	Ser	Asn	Arg	Ala	His	Gln	Gly	Lys	His	Glu
				85					90					95	

Thr Tyr Gly His Lys Thr Pro Tyr
100

<210> 3
 <211> 18
 <212> PRT
 <213> Homo sapiens

<400> 3

Thr	Glu	Val	Ser	His	His	Ile	Ser	Arg	Arg	Leu	Leu	Glu	Arg	Val	Asn
1				5					10					15	

Met Cys

<210> 4
 <211> 16
 <212> PRT
 <213> Homo sapiens

<400> 4

Lys	Asn	Gly	Lys	Gly	Asn	Val	Cys	His	Arg	Lys	Lys	His	His	Gly	Lys
				5					10					15	

<210> 5
 <211> 19
 <212> PRT
 <213> Homo sapiens

<400> 5

Asn	Ser	Asn	Arg	Ala	His	Gln	Gly	Lys	His	Glu	Thr	Tyr	Gly	His	Lys
1				5					10					15	

Thr Pro Tyr

<210> 6
 <211> 3117
 <212> DNA
 <213> Homo sapiens

<220>

<221> unsure
 <222> (1)..(3117)
 <223> n at any position in the sequence represents a or g or c or t/u

<220>
 <221> unsure
 <222> (1)..(3117)
 <223> y at any position in the sequence represents t/u or c

<220>
 <221> unsure
 <222> (1)..(3117)
 <223> m at any position in the sequence represents a or c

<220>
 <221> unsure
 <222> (1)..(3117)
 <223> k at any position in the sequence represents g or t/u

<220>
 <221> unsure
 <222> (1)..(3117)
 <223> s at any position in the sequence represents g or c

<220>
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 <222> (1)..(3117)
 <223> w at any position in the sequence represents a or t/u

<220>
 <221> unsure
 <222> (1)..(3117)
 <223> r at any position in the sequence represents g or a

<400> 6

aacatcctca cttgtgttgc tgtcagtgcc tgtanggcag gcaggaatgc agcagagagg	60
actcgccatc gtggccttgg ctgtctgtgc ggccctacat gcctcagaag ccatacttcc	120
cattgcctcc agctgttgca cggagggttc acatcatatt tccagaaggc tcctggaaag	180
agtgaatatg tgtcgcatcc agagagctga tggggattgt gacttggtg ctgtcatcct	240

tcatgtcaag	cgcnagaagaa	tctgtgtcag	cccgcacaac	catactgtta	agcagtggat	300
gaaagtgcaa	gctgccaaana	aaaatggtaa	aggaaatggt	tgccacagga	agaaacacca	360
tggcaagagg	aacagtaaca	gggcacatca	ggggaaacac	gaaacatacg	gccataaaac	420
tccttattag	agaatctaca	gataaatcta	cagagacaat	cccccaagtg	gacttggcca	480
tgattgggtg	taagttttatc	atctgaattc	tccttattgt	agacaacaga	acaaaacaaa	540
atattgggtt	ttaaaaaatg	aacaattgtg	ccgtatgcaa	atgtacccaa	taatatactc	600
cactggaaaa	tgaaatgaaa	aaannatact	ggctgggtat	ggtgggtccc	ccctttttatc	660
ccannnnctt	cgaggaggcag	aggcaggagg	atcacttgag	accaggantt	ngagacnagc	720
tnggggcaaa	anagcaanga	cntcatttnt	acaaacnaaa	aaaaannttg	gcccggcntg	780
gtagnacttg	cntataatcc	cagcnacatg	ggaggtngag	gtgggaggat	cacttgagtc	840
tggnngagtt	ngaggtngca	gtgagcagcn	tgggtgacag	aatgnagacc	ntgtctctaa	900
aaataataat	aataatgata	gtgtatatct	tcatataata	ttttaagnag	gagcatatag	960
atataacttn	ctcccaactt	tttaattata	gttttccaaa	cttacagaga	agttaaaaga	1020
atgggtacaat	gaacatctat	atatctttca	ccacaatatt	aatcattggt	aatattgtgc	1080
cacatttgct	ttctctctcc	tctcttggtg	ggggttncaa	tataaaatat	tataactttt	1140
aaaatatatc	ttgttttgct	aaccattgga	aaataagttg	caaaaatcat	gacacttcac	1200
ccctagtttc	ttttnggtgt	tataacttga	cataccctaa	aataaagaca	tttttctaca	1260
taatcacctt	atcagtttta	tacctaataa	attaataatt	tcatctaata	tattccatat	1320
tcaaattttc	ccaactattt	agagagcatt	ttatgtagtt	tttttttcac	tccagtaatc	1380
aatcaaggtn	gacatacata	ttgcaaataa	ttgttatatt	tctttaatat	ctttcaatct	1440
aagaaagttc	ctctgtcttt	tttttttaat	ttttaaaatt	attttggtga	gggaggggtc	1500
tgctgtgtct	tccaggctgg	agtgcagtgg	cacaattttg	attttggtc	actgaagcct	1560
caactttagg	gctcaagcaa	tcctcccacc	tcagcctncc	cgagtatctg	ggatcaaggt	1620
gcatacccac	cacacctggc	taattttgtt	tattttttgt	agagacaggg	tctcactatg	1680
ttgcccaggt	tgatctcaaa	ctcctgggct	caagcgatcc	tcccacctta	gcctcccaaa	1740

gtactgggat	tataggtgtg	agccacagtg	cctggcctaa	ttattttctt	gtgatcaaat	1800
tcagggttaa	tgtttttggt	taagaatttc	ctacgtgaat	tcgtgtactt	attttgtcat	1860
ttagagttca	taaatattag	ggtttatttt	ctaaatagaa	tagtttaaac	taaatataac	1920
ttcaaaacgt	ctagtttgag	tagctaccgt	tgtttggtt	gaaattttct	gatactgaaa	1980
agaacaaaaa	gcctgccttt	ctgcccanaa	csnnttgcyt	ccccagtna	gttcttgng	2040
cagnactagt	tagggnccca	gagttnggcc	ttnngkggtg	tgattttang	ytctgcctaa	2100
acaaggngcn	wacatytttt	agctcctatt	ccaccyttct	namamgtttt	tgttgtkggt	2160
tgnttggttt	tttkgagaca	grrtntnayt	ctgtttgccc	argctggart	tgcaagtggca	2220
caatytnngy	tncattgcaa	cytcngcytc	cssgccgttc	aaktgatyyt	cttgcytcag	2280
cytccccaa	taantgatat	tacaggngcc	cagccaccam	accccgntga	wttttgtatt	2340
tttartarar	amrgggtttt	cccgcnttgg	cngggctggt	ctcnaantcc	ttgamctcna	2400
ktgaaccacc	cgctgtgcc	ycccaaantg	ctggaattac	cancgttgan	ccaccatgcc	2460
gggcycacac	gtttgarttt	ganaccattg	tnccattcct	cttttggcct	yttttttntc	2520
catagnngct	tcaagataga	tangtaagrg	cccagtagtn	gttcwtarga	agcnmatagr	2580
rancrggar	cantttnatc	aggtgggcag	gtgtccnngg	cytccttgc	ggytnntccc	2640
aagcgggtgt	gttgccarga	nktnttggar	gtgataatgg	gananaccag	naggcmctga	2700
gtyncnntag	gttnaaatgc	caccaaact	ggcctttggc	ctaataatccy	ycnttgamta	2760
nttarcat	awtttattwa	tttnccctgac	atttntgcma	ncctttgtwt	ttntatttcc	2820
nctntatara	wgargaaatt	tgaggntytt	araggtaaaa	tganttgcnc	nrgtnnacmc	2880
aggaagtggc	nraranaanc	tttttanatn	mgaaaaaatt	aataaaatat	aatatgagag	2940
taacttaaaa	tattaataaa	ccacaatttt	aaattaatta	accgtgataa	ccaacattaa	3000
taaaagttaa	gataccaaaa	cactgggtgn	taattttttt	aactaacaan	ttgaattatt	3060
ttccatttta	aattaattaa	ccgtgataac	caacattaat	aaaagttaag	ataccgn	3117

<210> 7
 <211> 381
 <212> NA

<213> Homo sapiens

<220>

<221> unsure

<222> (208)..(208)

<223> n may represent a or g or c or t/u

<220>

<221> unsure

<222> (273)..(273)

<223> n may represent a or g or c or t/u

<400> 7

atgcagcaga gaggactcgc catcgtggcc ttggctgtct gtgcggccct acatgcctca 60

gaagccatac ttcccattgc ctccagctgt tgcacggagg ttccacatca tatttccaga 120

aggctcctgg aaagagtgaa tatgtgtcgc atccagagag ctgatgggga ttgtgacttg 180

gctgctgtca tccttcatgt caagcgcnga agaatctgtg tcagcccgca caaccatact 240

gttaagcagt ggatgaaagt gcaagctgcc aanaaaaatg gtaaaggaaa tgtttgccac 300

aggaagaaac accatggcaa gaggaacagt aacagggcac atcaggggaa acacgaaaca 360

tacggccata aaactcctta t 381

<210> 8

<211> 104

<212> DNA

<213> Homo sapiens

<400> 8

acacgaattc acgtaggaaa ttcttaacca aaaacattaa acctgaattt gatcacaaga 60

aaataattag gccaggcact gtggctcaca cctataatcc cagt 104

<210> 9

<211> 25

<212> DNA

<213> Homo sapiens

<400> 9

gaattcacgt aggaaattct taacc 25

<210> 10
<211> 22
<212> DNA
<213> Homo sapiens

<400> 10
actgggatta taggtgtgag cc

22

<210> 11
<211> 311
<212> DNA
<213> Homo sapiens

<220>
<221> unsure
<222> (101)..(101)
<223> n may be a or g or c or t/u

<220>
<221> unsure
<222> (162)..(162)
<223> n may be a or g or c or t/u

<400> 11
ggagagagcc gtatgttttcg tgtttcccct gatgtgccct gttactgttc ctcttgccat 60
gggtgtttcct cctgtggcaa acatttcctt taccattttt nttggcagct tgcactttca 120
tccactgctt aacagtatgg ttgtgcgggc tgacacagat tnttctgcgc ttgacatgaa 180
gggatgacagc agccaagtca caatccccat cagctctctg gatgcgacac atattcactc 240
tttccaggag ccttctggaa atatgatgtg aaacctccgt gcaacagctg gaggcaatgg 300
gaagtatggc t 311

<210> 12
<211> 20
<212> DNA
<213> Artificial sequence

<220>
<223> Sequencing primer T7

<400> 12
taatacgact cactataggg

20

<210> 13
<211> 18
<212> DNA
<213> Artificial sequence

<220>
<223> pCR3.1 Reverse Primer

<400> 13
tagaaggcac agtcgagg

18

<210> 14
<211> 22
<212> DNA
<213> Artificial sequence

<220>
<223> Gene specific primer (24R)

<400> 14
actgggatta taggtgtgag cc

22

<210> 15
<211> 24
<212> DNA
<213> Artificial sequence

<220>
<223> Gene specific primer (24R2)

<400> 15
caaattcagg tttaatgttt ttgg

24

<210> 16
<211> 21
<212> DNA
<213> Artificial sequence

<220>
<223> Gene specific primer (F4)

<400> 16
ctcaaacgtg tgagcccggc a

21

<210> 17

<211> 25
<212> DNA
<213> Artificial sequence

<220>
<223> Gene specific primer (F3)

<400> 17
gctactcaaa ctagacgttt tgaag

25

<210> 18
<211> 24
<212> DNA
<213> Artificial sequence

<220>
<223> primers F8

<400> 18
ccgtatgttt cgtgtttccc ctga

24

<210> 19
<211> 24
<212> DNA
<213> Artificial sequence

<220>
<223> Primer R5

<400> 19
agccatactt cccattgcct ccag

24

<210> 20
<211> 150
<212> PRT
<213> Homo sapiens

<400> 20

Met Asn Leu Trp Leu Leu Ala Cys Leu Val Ala Gly Phe Leu Gly Ala
1 5 10 15

Trp Ala Pro Ala Val His Thr Gln Gly Val Phe Glu Asp Cys Cys Leu
20 25 30

Ala Tyr His Tyr Pro Ile Gly Trp Ala Val Leu Arg Arg Ala Trp Thr
35 40 45

Tyr Arg Ile Gln Glu Val Ser Gly Ser Cys Asn Leu Pro Ala Ala Ile
50 55 60

Phe Tyr Leu Pro Lys Arg His Arg Lys Val Cys Gly Asn Pro Lys Ser
65 70 75 80

Arg Glu Val Gln Arg Ala Met Lys Leu Leu Asp Ala Arg Asn Lys Val
85 90 95

Phe Ala Lys Leu His His Asn Met Gln Thr Phe Gln Ala Gly Pro His
100 105 110

Ala Val Lys Lys Leu Ser Ser Gly Asn Ser Lys Leu Ser Ser Ser Lys
115 120 125

Phe Ser Asn Pro Ile Ser Ser Ser Lys Arg Asn Val Ser Leu Leu Ile
130 135 140

Ser Ala Asn Ser Gly Leu
145 150

<210> 21

<211> 95

<212> PRT

<213> Homo sapiens

<400> 21

Met Cys Cys Thr Lys Ser Leu Leu Leu Ala Ala Leu Met Ser Val Leu
5 10 15

Leu Leu His Leu Cys Gly Glu Ser Glu Ala Ser Asn Phe Asp Cys Cys
20 25 30

Leu Gly Tyr Thr Asp Arg Ile Leu His Pro Lys Phe Ile Val Gly Phe
35 40 45

Thr Arg Gln Leu Ala Asn Glu Gly Cys Asp Ile Asn Ala Ile Ile Phe
50 55 60

His Thr Lys Lys Lys Leu Ser Val Cys Ala Asn Pro Lys Gln Thr Trp
65 70 75 80

Val Lys Tyr Ile Val Arg Leu Leu Ser Lys Lys Val Lys Asn Met
85 90 95

<210> 22

<211> 94

<212> PRT

<213> Homo sapiens

<400> 22

Met Ala Pro Leu Lys Met Leu Ala Leu Val Thr Leu Leu Leu Gly Ala
1 5 10 15

Ser Leu Gln His Ile His Ala Ala Arg Gly Thr Asn Val Gly Arg Glu
20 25 30

Cys Cys Leu Glu Tyr Phe Lys Gly Ala Ile Pro Leu Arg Lys Leu Lys
35 40 45

Thr Trp Tyr Gln Thr Ser Glu Asp Cys Ser Arg Asp Ala Ile Val Phe
50 55 60

Val Thr Val Gln Gly Arg Ala Ile Cys Ser Asp Pro Asn Asn Gln Arg
65 70 75 80

Val Lys Asn Ala Val Lys Tyr Leu Gln Ser Leu Glu Arg Ser
85 90

210> 23
211> 96
212> PRT
213> Homo sapiens

400> 23

Met Gln Ile Ile Thr Thr Ala Leu Val Cys Leu Leu Leu Ala Gly Met
5 10 15

Trp Pro Glu Asp Val Asp Ser Lys Ser Met Gln Val Pro Phe Ser Arg
20 25 30

Cys Cys Phe Ser Phe Ala Glu Gln Glu Ile Pro Leu Arg Ala Ile Leu
35 40 45

Cys Tyr Arg Asn Thr Ser Ser Ile Cys Ser Asn Glu Gly Leu Ile Phe
50 55 60

Lys Leu Lys Arg Gly Lys Glu Ala Cys Ala Leu Asp Thr Val Gly Trp
65 70 75 80

Val Gln Arg His Arg Lys Met Leu Arg His Cys Pro Ser Lys Arg Lys
85 90 95

<210> 24
<211> 77
<212> PRT
<213> Homo sapiens

<400> 24

Ala Gln Pro Asp Ser Val Ser Ile Pro Ile Thr Cys Cys Phe Asn Val
1 5 10 15

Ile Asn Arg Lys Ile Pro Ile Gln Arg Leu Glu Ser Tyr Thr Arg Ile
20 25 30

Thr Asn Ile Gln Cys Pro Lys Glu Ala Val Ile Phe Lys Thr Lys Arg
35 40 45

Gly Lys Glu Val Cys Ala Asp Pro Lys Glu Arg Trp Val Arg Asp Ser
50 55 60

Met Lys His Leu Asp Gln Ile Phe Gln Asn Leu Lys Pro
65 70 75

<210> 25

<211> 98

<212> PRT

<213> Homo sapiens

<400> 25

Met Lys Val Ser Ala Val Leu Leu Cys Leu Leu Leu Met Thr Ala Ala
5 10 15

Phe Asn Pro Gln Gly Leu Ala Gln Pro Asp Ala Leu Asn Val Pro Ser
20 25 30

Thr Cys Cys Phe Thr Phe Ser Ser Lys Lys Ile Ser Leu Gln Arg Leu
35 40 45

Lys Ser Tyr Val Ile Thr Thr Ser Arg Cys Pro Gln Lys Ala Val Ile
50 55 60

Phe Arg Thr Lys Leu Gly Lys Glu Ile Cys Ala Asp Pro Lys Glu Lys
65 70 75 80

Trp Val Gln Asn Tyr Met Lys His Leu Gly Arg Lys Ala His Thr Leu
85 90 95

Lys Thr

<210> 26

<211> 97

<212> PRT

<213> Homo sapiens

<400> 26

Met	Lys	Val	Ser	Ala	Ala	Leu	Leu	Trp	Leu	Leu	Leu	Ile	Ala	Ala	Ala
1				5					10					15	
Phe	Ser	Pro	Gln	Gly	Leu	Ala	Gly	Pro	Ala	Ser	Val	Pro	Thr	Thr	Cys
			20					25					30		
Cys	Phe	Asn	Leu	Ala	Asn	Arg	Lys	Ile	Pro	Leu	Gln	Arg	Leu	Glu	Ser
		35					40					45			
Tyr	Arg	Arg	Ile	Thr	Ser	Gly	Lys	Cys	Pro	Gln	Lys	Ala	Val	Ile	Phe
	50					55					60				
Lys	Thr	Lys	Leu	Ala	Lys	Asp	Ile	Cys	Ala	Asp	Pro	Lys	Lys	Lys	Trp
65					70					75					80
Val	Gln	Asp	Ser	Met	Lys	Tyr	Leu	Asp	Gln	Lys	Ser	Pro	Thr	Pro	Lys
				85					90					95	

Pro

<210> 27

<211> 99

<212> PRT

<213> Homo sapiens

<400> 27

Met	Lys	Ala	Ser	Ala	Ala	Leu	Leu	Cys	Leu	Leu	Leu	Thr	Ala	Ala	Ala
				5					10					15	
Phe	Ser	Pro	Gln	Gly	Leu	Ala	Gln	Pro	Val	Gly	Ile	Asn	Thr	Ser	Thr
			20					25					30		
Thr	Cys	Cys	Tyr	Arg	Phe	Ile	Asn	Lys	Lys	Ile	Pro	Lys	Gln	Arg	Leu
		35					40					45			
Glu	Ser	Tyr	Arg	Arg	Thr	Thr	Ser	Ser	His	Cys	Pro	Arg	Glu	Ala	Val
	50					55					60				
Ile	Phe	Lys	Thr	Lys	Leu	Asp	Lys	Glu	Asp	Cys	Ala	Asp	Pro	Thr	Gln
65					70					75					80
Lys	Trp	Val	Gln	Asp	Pro	Met	Lys	His	Leu	Asp	Lys	Lys	Thr	Gln	Thr
				85					90					95	

Pro Lys Leu

<210> 28
 <211> 99
 <212> PRT
 <213> Homo sapiens

<400> 28

Met	Lys	Val	Ser	Ala	Ala	Leu	Leu	Cys	Leu	Leu	Leu	Thr	Ala	Ala	Ala
1				5					10					15	
Phe	Ile	Pro	Gln	Gly	Leu	Ala	Gln	Pro	Asp	Ala	Ile	Asn	Ala	Pro	Val
			20					25					30		
Thr	Cys	Cys	Tyr	Asn	Phe	Thr	Asn	Arg	Lys	Ile	Ser	Val	Gln	Arg	Leu
		35					40					45			
Ala	Ser	Tyr	Arg	Arg	Ile	Thr	Ser	Ser	Lys	Cys	Pro	Lys	Glu	Ala	Val
	50					55					60				
Ile	Phe	Lys	Thr	Ile	Val	Ala	Lys	Glu	Asp	Cys	Ala	Asp	Pro	Lys	Gln
65					70					75					80
Lys	Trp	Val	Gln	Asp	Ser	Met	Asp	His	Leu	Asp	Lys	Gln	Thr	Gln	Thr
				85					90					95	
Pro	Lys	Thr													

<210> 29
 <211> 91
 <212> PRT
 <213> Homo sapiens

<400> 29

Met	Lys	Val	Ser	Ala	Ala	Arg	Leu	Ala	Val	Ile	Leu	Ile	Ala	Thr	Ala
1				5					10					15	
Leu	Cys	Ala	Pro	Ala	Ser	Ala	Ser	Pro	Tyr	Ser	Ser	Asp	Thr	Thr	Pro
			20					25					30		
Cys	Cys	Phe	Ala	Tyr	Ile	Ala	Arg	Pro	Leu	Pro	Arg	Ala	His	Ile	Lys
		35					40					45			
Glu	Tyr	Phe	Tyr	Thr	Ser	Gly	Lys	Cys	Ser	Asn	Pro	Ala	Val	Val	Phe
	50					55					60				
Val	Thr	Arg	Lys	Asn	Arg	Gln	Val	Cys	Ala	Asn	Pro	Glu	Lys	Lys	Trp
65					70					75					80

Val Arg Glu Tyr Ile Asn Ser Leu Glu Met Ser
85 90

<210> 30
<211> 93
<212> PRT
<213> Homo sapiens

<400> 30

Met Lys Ile Ser Val Ala Ala Ile Pro Phe Phe Leu Leu Ile Thr Ile
1 5 10 15

Ala Leu Gly Thr Lys Thr Glu Ser Ser Ser Arg Gly Pro Tyr His Pro
20 25 30

Ser Glu Cys Cys Phe Thr Tyr Thr Thr Tyr Lys Ile Pro Arg Gln Arg
35 40 45

Ile Met Asp Tyr Tyr Glu Thr Asn Ser Gln Cys Ser Lys Pro Gly Ile
50 55 60

Val Phe Ile Thr Lys Arg Gly His Ser Val Cys Thr Asn Pro Ser Asp
65 70 75 80

Lys Trp Val Gln Asp Tyr Ile Lys Asp Met Lys Glu Asn
85 90

<210> 31
<211> 92
<212> PRT
<213> Homo sapiens

<400> 31

Met Lys Leu Cys Val Thr Val Leu Ser Leu Leu Met Leu Val Ala Ala
1 5 10 15

Phe Cys Ser Pro Ala Leu Ser Ala Pro Met Gly Ser Asp Pro Pro Thr
20 25 30

Ala Cys Cys Phe Ser Tyr Thr Ala Arg Lys Leu Pro Arg Asn Phe Val
35 40 45

Val Asp Tyr Tyr Glu Thr Ser Ser Leu Cys Ser Gln Pro Ala Val Val
50 55 60

Phe Gln Thr Lys Arg Ser Lys Gln Val Cys Ala Asp Pro Ser Glu Ser
65 70 75 80

Trp Val Gln Glu Tyr Val Tyr Asp Leu Glu Leu Asn
85 90

<210> 32
<211> 93
<212> PRT
<213> Homo sapiens

<400> 32

Met Gln Val Ser Thr Ala Ala Leu Ala Val Leu Leu Cys Thr Met Ala
1 5 10 15

Leu Cys Asn Gln Val Leu Ser Ala Pro Leu Ala Ala Asp Thr Pro Thr
20 25 30

Ala Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe Ile
35 40 45

Ala Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Ser Val Ile
50 55 60

Phe Leu Thr Lys Arg Gly Arg Gln Val Cys Ala Asp Pro Ser Glu Glu
65 70 75 80

Trp Val Gln Lys Tyr Val Ser Asp Leu Glu Leu Ser Ala
85 90

<210> 33
<211> 92
<212> PRT
<213> Homo sapiens
<400> 33

Met Gln Val Ser Thr Ala Ala Leu Ala Val Leu Leu Cys Thr Met Ala
1 5 10 15

Leu Cys Asn Gln Phe Ser Ala Ser Leu Ala Ala Asp Thr Pro Thr Ala
20 25 30

Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe Ile Ala
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Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Gly Val Ile Phe
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35 40 45

Glu Thr Ser Pro Gln Cys Pro Lys Pro Gly Val Ile Leu Leu Thr Lys
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